Listing of the Claims

This listing of claims will replace all prior versions, and listings of claims in the application.

- 1. (Currently amended) A method for identifying a compound that inhibits sister chromatid separation comprising inhibiting the proteolytic activity of separase, wherein an active separase in the form of a full-length separase upon activation in the presence of securin is incubated in the presence of a separase substrate with a test compound, and wherein the inhibiting effect of the test compound on the proteolytic activity of the active separase is determined; wherein said separase substrate is a peptide comprising an amino acid sequence EXXR, wherein X is any amino acid, and the separase substrate is capable of being cleaved by the active separase.
 - 2. (Original) The method of claim 1, wherein the active separase is human.
- 3. (Previously presented) The method of claim 1, wherein the active separase has been obtained by activation of the full-length separase in a mitotic cell extract in the presence of securin.
- 4. (Original) The method of claim 3, wherein the mitotic cell extract has been obtained from Xenopus laevis eggs.

- 5. (Previously presented) The method of claim 1, wherein the separase substrate is a peptide comprising a fluorogenic group, wherein processing of the peptide results in a change in fluorescence, and wherein the change in fluorescence is correlated with the separase activity.
- 6. (Previously presented) The method of claim 5, wherein the separase substrate is a peptide comprising an amino acid sequence DREIMR (SEQ ID NO:9), SFEILR (SEQ ID NO:11) or EWELLR (SEQ ID NO:12).

7 - 10. (Cancelled)

- chromatid separation comprising inhibiting the proteolytic activity of separase, wherein an active separase in the form of one or more separase fragments, optionally upon activation of a full-length separase in the presence of securin, is incubated in the presence of a separase substrate with a test compound, and wherein the inhibiting effect of the test compound on the proteolytic activity of the active separase is determined; wherein said separase substrate is a peptide comprising an amino acid sequence EXXR, wherein X is any amino acid, and the separase substrate is capable of being cleaved by the active separase.
- 12. (Previously presented) The method of claim 11, wherein the active separase is human.

- 13. (Previously presented) The method of claim 11, wherein the active separase has been obtained by activation of one or more separase fragments in a mitotic cell extract in the presence of securin.
- 14. (Previously presented) The method of claim 13, wherein the mitotic cell extract has been obtained from Xenopus laevis eggs.
- 15. (Previously presented) The method of claim 11, wherein the separase substrate is a peptide comprising a fluorogenic group, wherein processing of the peptide results in a change in fluorescence, and wherein the change in fluorescence is correlated with the separase activity.
- 16. (Previously presented) The method of claim 15, wherein the separase substrate is a peptide comprising an amino acid sequence DREIMR (SEQ ID NO:9), SFEILR (SEQ ID NO:11) or EWELLR (SEQ ID NO:12).